

IN THE SPECIFICATION

On page 14, the paragraph starting on line 11, please replace with the following paragraph:

Analysis of the DNA encoding z219c polypeptide (SEQ ID NO:1) revealed an open reading frame encoding 223 amino acids (SEQ ID NO:2) comprising a predicted signal peptide of 21 amino acid residues (residue 1 (Met) to residue 21 (Met) of SEQ ID NO:2), and a mature polypeptide of 194 amino acids (residue 22 (Phe) to residue 223 (Phe) of SEQ ID NO:2). Multiple alignment of z219c with other members of the human 2-19 protein family revealed the following 3 regions of conserved amino acids (see Figure):

IN THE CLAIMS

Please cancel claims, 1, 4, 6, 7, 8, 9, 23, and 25 without prejudice to the prosecution thereof in a continuing application.

Please amend the following claims:

B² 3. (Amended) An isolated polynucleotide sequence that comprises nucleotide 1 to nucleotide 669 or nucleotide 64 to nucleotide 699 of SEQ ID NO:8.

B³ 5. (Twice Amended) An isolated polynucleotide according to claim 3, wherein the polynucleotide consists of nucleotide 1 to nucleotide 699 or nucleotide 64 to nucleotide 699 of SEQ ID NO:8.

B⁴ 22. (Amended) A vector comprising the following operably linked elements:
a transcription promoter;
a DNA segment comprising a polynucleotide selected from the group consisting of: